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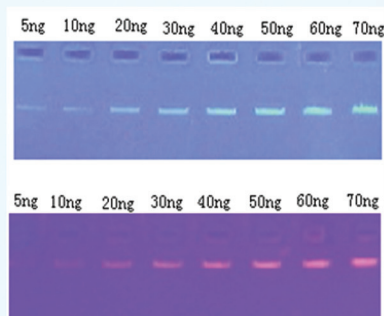
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GoodView™ Nucleic Acid Stain

—An alternative to EB

GoodView™ is a new nucleic acid stain, an alternative to the traditional ethidium bromide (EB) stain for detecting nucleic acid in agarose gels. It emits green fluorescence when bound to DNA or RNA. This new stain has two fluorescence excitation maxima when bound to nucleic acid, one centered at 268 nm and another at 294 nm. In addition, it has one visible excitation at 491 nm. The Fluorescence emission of GoodView™ bound to DNA is centered at 530 nm.

Comparative sensitivity test of GV and EB



Sensitivity test result of
GV at UV 300nm.

Sensitivity test result of
EB at UV 300nm.

The result of electrophoresis demonstrates GV is almost as sensitive as EB.

The Test Report from Institute for Environmental Health and Related Product Safety of Chinese Center for Disease Control and Prevention concludes that:

- ◆ Acute Oral Toxicity Test: GoodView™ Nucleic Acid Stain belongs to nontoxic.
- ◆ Mouse Marrow Chromophilous Erythrocyte Micronucleus Test: Negative. There is no significant difference in the incidence of micronuclei between test and control groups.
- ◆ Ames Test: Negative. No mutagenicity was observed.
- ◆ In Vitro Mammalian Cell Chromosome Aberration Test: Negative. No increasing aberration rate was observed.

The test report is available upon request.

Echo® Acoustic LIQUID HANDLING
for GENOMICS



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Echo® Liquid Handlers enable library preparation in low microliter volumes for a range of sequencing methods. Dramatically reduce reagent costs, conserve samples, and eliminate steps – all while improving library quality.

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- ▶ 100-fold reduction of library prep reaction volumes
- ▶ 30-fold reduction of sample pooling turnaround time
- ▶ Increased sample throughput
- ▶ Automation of workflow to easily prepare thousands of samples
- ▶ Improved accuracy of results

Comparison of Liquid Handling Methods*

	Manual Pipetting	Echo® Liquid Handler
Amount of DNA	50 ng	0.06 – 2.0 ng
DNA volume (Rxn)	25 µL	200 nL
Library prep volume (Rxn)	25 µL	300 nL
Total volume	50 µL	0.5 µL
Reactions per kit	96	9600
Cost per reaction	\$72.91	\$0.73

For more information, visit www.labcyte.com/sequencing.

* Low-Cost, High-Throughput Sequencing of DNA Assemblies Using a Highly Multiplexed Nextera Process. Shapland et al. ACS Synth. Biol., 2015

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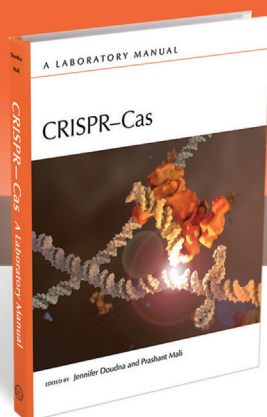
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CRISPR-Cas

A Laboratory Manual



The essential guide to CRISPR-Cas

Edited by Jennifer Doudna, *University of California, Berkeley*;
Prashant Mali, *University of California, San Diego*

The development of CRISPR-Cas technology is revolutionizing biology. Based on machinery bacteria use to target foreign nucleic acids, these powerful techniques allow investigators to edit nucleic acids and modulate gene expression more rapidly and accurately than ever before.

Featuring contributions from leading figures in the CRISPR-Cas field, this laboratory manual presents a state-of-the-art guide to the technology. It includes step-by-step protocols for applying CRISPR-Cas-based techniques in various systems, including yeast, zebrafish, *Drosophila*, mice, and cultured cells (e.g., human pluripotent stem cells). The contributors cover web-based tools and approaches for designing guide RNAs that precisely target genes of interest, methods for preparing and delivering CRISPR-Cas reagents into cells, and ways to screen for cells that harbor the desired genetic changes. Strategies for optimizing CRISPR-Cas in each system—especially for minimizing off-target effects—are also provided.

Authors also describe other applications of the CRISPR-Cas system, including its use for regulating genome activation and repression, and discuss the development of next-generation CRISPR-Cas tools. The book is thus an essential laboratory resource for all cell, molecular, and developmental biologists, as well as biochemists, geneticists, and all who seek to expand their biotechnology toolkits.

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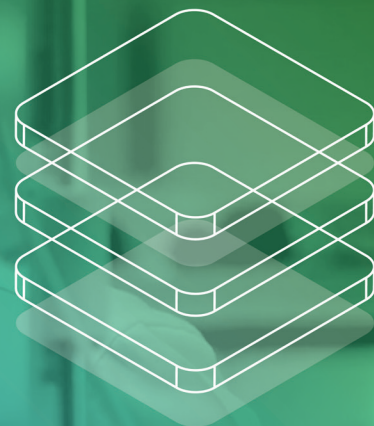
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The AACR Science of Cancer Health Disparities conferences advance the understanding of, and ultimately help to eliminate, the disparities that represent a major public health problem in our country. Reflecting this transdisciplinary field, professionals from academia, industry, government, and the community are brought together to promote the exchange of novel ideas, discuss the latest findings in the field, and stimulate the development of new research on cancer health disparities.

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Conference Cochairs: Carlos L. Arteaga,
Carlos Gil M. Ferreira, and Gabriel A. Rabinovich
September 27-29, 2018 | São Paulo, Brazil

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Conference Cochairs: Anil K. Rustgi, Johanna Bendell,
Hans Clevers, Christina Curtis, and Owen Sansom
September 27-30, 2018 | Washington, DC

Metabolism and Cancer

Conference Cochairs: Ralph J. Deberardinis, Tak W. Mak,
Joshua D. Rabinowitz, and M. Celeste Simon
September 28-October 1, 2018 | New York, NY

Fourth CRI-CIMT-EATI-AACR International Cancer Immunotherapy Conference: Translating Science into Survival

Conference Cochairs: Nina Bhardwaj, Christoph
Huber, Elizabeth M. Jaffee, and Guido Kroemer
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EACR-AACR-ISCR Conference: The Cutting Edge of Contemporary Cancer Research

Conference Cochairs: Richard M. Marais,
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Laura Fejerman, Scarlett Lin Gomez,
Augusto C. Ochoa, and Brian M. Rivers
November 2-5, 2018 | New Orleans, LA

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Scientific Committee Cochairs: Charles Swanton,
James L. Gulley, and Antoni Ribas
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Tumor Immunology and Immunotherapy

Conference Cochairs: James P. Allison,
Lisa M. Coussens, Ira Mellman, and Drew M. Pardoll
November 27-30, 2018 | Miami Beach, FL

Innovation and Biomarkers in Cancer Drug Development: A Joint Meeting Presented By EORTC, NCI, EMA, and AACR

Organizing Committee Chair: Denis A. Lacombe
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Targeting PI3K/mTOR Signaling

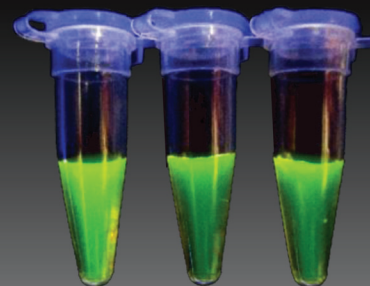
Conference Cochairs: Lewis C. Cantley,
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- Fluorescence emitted by Eprobe can be detected using a filter for SYBR® Green I. *SYBR is a registered trademark of Molecular Probes, Inc.

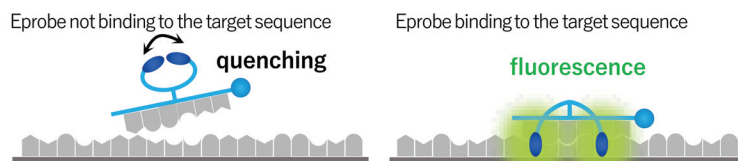


Figure 1. Fluorescence emission mechanism of Eprobe

High resolution SNP detection with Eprobe

- Melting curve analysis after asymmetric PCR with Eprobe can detect genotype of SNP (Fig.2).
- Increased T_m of the Eprobe enables a shorter probe design and a clearer distinction of single nucleotide substitution.
- Predesigned Eprobes targeting SNP for ADH1B (rs1229984), ADRB2 (rs1042713), ALDH2 (rs671), FTO (rs9939609), UCP1 (rs1800592) and others are available.

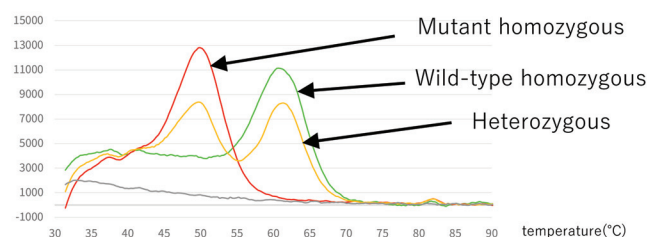


Figure 2. Predesigned Eprobe for IL28B (rs8099917).

Highly sensitive somatic mutation detection

- Highly sensitive detection of somatic mutations (down to 0.1%) can be achieved (Fig.3) by suppression of PCR amplification of wild-type alleles by Eprobe (PCR clamping).

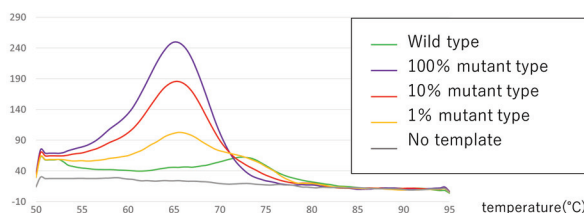


Figure 3. Predesigned Eprobe for G12D in the KRAS gene.

Pricing and ordering information

Product	Fluorophore	Quantity	List price
Eprobe Modification: 3' Spacer C3.	Thiazole Orange	1.5 nmol	¥19,000 ¥38,000
		3.0 nmol	¥30,000 ¥60,000
		5.0 nmol	¥45,000 ¥90,000
		10.0 nmol	¥70,000 ¥140,000
	Thiazole Pink	1.5 nmol	¥45,000
		3.0 nmol	¥70,000
		5.0 nmol	¥110,000
		10.0 nmol	¥170,000

- Excitation/Emission wave length (nm): Thiazole Orange: 510/530, Thiazole Pink: 570/590.
- Purification: HPLC, Shipping format: dry.
- Shipping charge: 11,000 JPY/ shipment.

Product	Fluorophore	Quantity	List price
Eprimer 3' unmodified: Extension from the 3' end is possible.	Thiazole Orange	1.5 nmol	¥19,000 ¥38,000
		3.0 nmol	¥30,000 ¥60,000
		5.0 nmol	¥45,000 ¥90,000
		10.0 nmol	¥70,000 ¥140,000
	Thiazole Pink	1.5 nmol	¥45,000
		3.0 nmol	¥70,000
		5.0 nmol	¥110,000
		10.0 nmol	¥170,000

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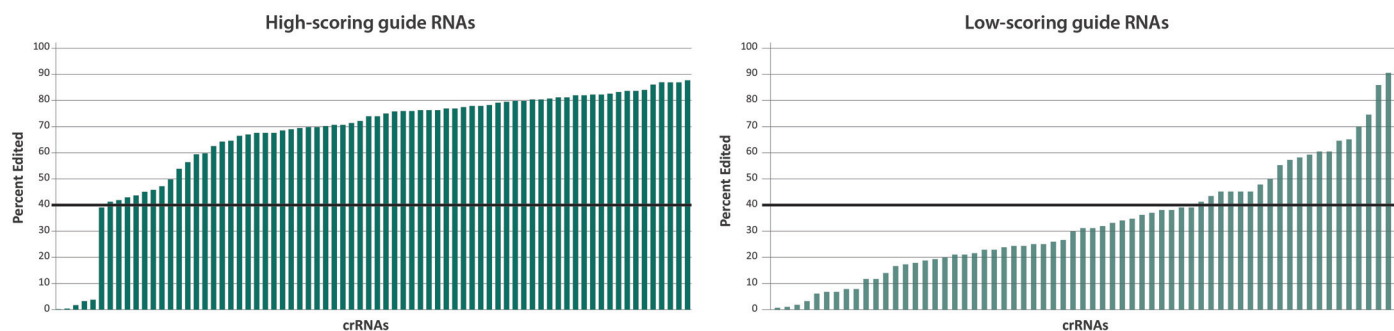


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CRISPR Guide RNAs | CRISPR Screening Libraries | Cas9 Nucleases

crRNAs that have high functionality scores show high editing efficiency



HEK293T-CAG-Cas9 cells were transfected with either high-scoring or low-scoring crRNAs (50 nM crRNA:tracrRNA) using DharmaFECT 1 transfection reagent (0.25 μ L/well) in 96-well format. Gene editing efficiencies were determined using next-generation sequencing. 93% of the top 10 high-scoring crRNAs targeting ten different genes have > 40% indel formation and only 33% of the 10 lowest scoring designs have > 40% indel formation.